SEQUENCE LISTING



<110> PANACCIO, Michael HASSE, Detlef

<120> THERAPEUTIC AND DIAGNOSTIC COMPOSITIONS

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<151> 1995-11-30

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<151> 1996-11-29

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<170> PatentIn Ver. 2.0

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att aca aaa gat ggt gta tct gtt gca aaa gaa att gaa ctt gaa gat 192 Ile Thr Lys Asp Gly Val Ser Val Ala Lys Glu Ile Glu Leu Glu Asp 50 55 60

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act agc gat att gct ggt gat gga act aca aca gca aca gtc ctt gca 288 Thr Ser Asp Ile Ala Gly Asp Gly Thr Thr Ala Thr Val Leu Ala

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gct c Ala G 145															480
atc a Ile I		_	_	_	-	_		_							528
gtt g Val G	_	-	_				_				_		-	-	 576
atg a Met L	_		_	_								_			624
gag a Glu L 210		_	-	-	_								-		 672
aaa a Lys L 225															720
gct a Ala L															768
gaa g Glu A			_			_	-		-		_		-		816
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ctt g Leu G 2															912
cgt g Arg G 305															960

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Lys Phe Glu Asn Met Gly Ala Gln Met Val Lys Glu Val Ala Pro Lys 65 70 75 80

Thr Ser Asp Ile Ala Gly Asp Gly Thr Thr Thr Ala Thr Val Leu Ala 85 90 95

Gln Ala Ile Tyr Arg Glu Gly Val Lys Leu Val Ala Ala Gly Arg Asn 100 105 110

Pro Met Ala Ile Lys Arg Gly Ile Asp Lys Ala Val Val Ala Val Thr 115 120 125

Lys Glu Leu Ser Asp Ile Thr Lys Pro Thr Arg Asp Gln Lys Glu Ile 130 135 140

Ala Gln Val Gly Thr Ile Ser Ala Asn Ser Asp Thr Thr Ile Gly Asn 145 150 155 160

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Arg G. 305	ly Iḷe	Lys	Leu	Glu 310	Asn	Val	Ser	Leu	Ser 315	Ser	Leu	Gly	Thr	Ala 320
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Inr A.	la Phe	Val 420	ALG		116	-1-	425		ч	P		430		
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Thr Asp Asp Gly Lys Leu Ile Pro Met Ala Val Lys Ala Gly Asp Thr
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Leu Ala Ile Gly Phe Thr Gly Ser Gln Gly Pro Asn Gln Ala Gly Met $35 \hspace{1cm} 40 \hspace{1cm} 45$

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Lys Thr Leu Asn Asp Leu Asp Ile Leu Leu Lys Asp Val Met Leu Thr 20 25 30

Ser Lys Lys His Glu Ser Arg Arg Leu Ala Glu Ser Val His Gln Asn

35 40 45

Ile Leu Thr His Leu Ile Gln Lys Asn Tyr Asn Thr His Asn Gly Gly 50 55 60

Ile Lys Ser Ala Pro Phe His Val Leu Ile Gly Pro Lys Ile Pro Ser 65 70 75 80

Ile Leu Val Glu Val Gly Tyr Cys Ser Asn Lys Ala Glu Ala Gln Arg 85 90 95

Leu Ala Ser Ser Asn Tyr Gln Lys Ala Leu Ile Glu Gly Leu Ala Lys
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20 25 30

Ala Ser Pro Asn Glu Ile Ser Phe Leu Ala Asn Ala Lys Tyr Ile His
35 40 45

Gln Leu Val Leu Ser Gln Ala Gly Ala Ile Ile Leu Ser Lys Glu Tyr 50 55 60

Ala Ser Arg Val Pro Arg Ala Leu Ile Ser Thr Glu Pro Tyr Arg Asp
65 70 75 80

Phe Gly Arg Val Leu Ser Leu Phe Ser Ile Pro Gln Gly Cys Phe Asp 85 90 95

Gly Ile Ser His Gln Ala Tyr Ile His Pro Thr Ala Gln Val Ser Lys 100 105 110

Thr Ala Thr Ile Tyr Pro Xaa Val Phe Ile Gly
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aaa gaa gac ttt ggg cgt cat att gtt aaa tta tgc tgg aaa ggt tct Lys Glu Asp Phe Gly Arg His Ile Val Lys Leu Cys Trp Lys Gly Ser 20 25 30	95													
tta tca aat atc ttt ttt tcc tat ggg gat atc ccg cac cca cct tat Leu Ser Asn Ile Phe Phe Ser Tyr Gly Asp Ile Pro His Pro Pro Tyr 35 40 45	143													
ata cat caa agt aat aag gtt cag gat aag gaa aga tat cnt acn gta Ile His Gln Ser Asn Lys Val Gln Asp Lys Glu Arg Tyr Xaa Xaa Val 50 55 60	191													
tac tot ata tta cat aan otg ggt tot gta gca gct cot aca gct gga Tyr Ser Ile Leu His Xaa Leu Gly Ser Val Ala Ala Pro Thr Ala Gly 65 70 75	239													
tta cnc ttt tct gaa act agc cgt nat aaa tta cac aaa nat ggt att Leu Xaa Phe Ser Glu Thr Ser Arg Xaa Lys Leu His Lys Xaa Gly Ile 80 85 90 95	287													
agt tgg gca taa atc cct ctt cac gtg gga tat gga aca ttc agt ccc Ser Trp Ala Ile Pro Leu His Val Gly Tyr Gly Thr Phe Ser Pro 100 105 110	335													
gtc ctc tgc aat gac atc cca aaa cat ctt atc cnt tct gag ttt gtt Val Leu Cys Asn Asp Ile Pro Lys His Leu Ile Xaa Ser Glu Phe Val 115 120 125	383													
cac ttt cct gaa act acn ttt tcc act ata tta aat gca cgg ttt gca His Phe Pro Glu Thr Xaa Phe Ser Thr Ile Leu Asn Ala Arg Phe Ala 130 135 140	431													
ngg gaa tac cta tgt tct gcc ata ggg gac cca ctg ttg tcc cca cca Xaa Glu Tyr Leu Cys Ser Ala Ile Gly Asp Pro Leu Leu Ser Pro Pro 145 150 155	479													
ttg gan ggg tgt tat ctt acc cct ttc gcc cgg ggt tcc cct ccc caa Leu Xaa Gly Cys Tyr Leu Thr Pro Phe Ala Arg Gly Ser Pro Pro Gln 160 165 170	527													
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<211> 98

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Ser Asn Ile Phe Phe Ser Tyr Gly Asp Ile Pro His Pro Pro Tyr Ile 35 40 45

His Gln Ser Asn Lys Val Gln Asp Lys Glu Arg Tyr Xaa Xaa Val Tyr 50 55 60

Ser Ile Leu His Xaa Leu Gly Ser Val Ala Ala Pro Thr Ala Gly Leu 65 70 75 80

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Trp Ala

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<213> Lawsonia intracellularis

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Asp Ile Pro Lys His Leu Ile Xaa Ser Glu Phe Val His Phe Pro Glu 20 25 30

Thr Xaa Phe Ser Thr Ile Leu Asn Ala Arg Phe Ala Xaa Glu Tyr Leu 35 40 45

ys Ser Ala Ile Gly Asp Pro Leu Leu Ser Pro Pro Leu Xaa Gly Cys 50 55 60

Tyr Leu Thr Pro Phe Ala Arg Gly Ser Pro Pro Gln Pro Tyr Ser Ile
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Xaa Phe Ser Ser Gln Ile

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tat ata gga gaa cat tca cca tta aaa cgt aat gta ant atg gaa gat
Tyr Ile Gly Glu His Ser Pro Leu Lys Arg Asn Val Xaa Met Glu Asp
                 20
                                     25
                                                         30
gta ggt aaa tot got gtt ttt tta got toa gac otn toa toa gga gta
                                                                  145
Val Gly Lys Ser Ala Val Phe Leu Ala Ser Asp Xaa Ser Ser Gly Val
             35
                                 40
acc ggt gaa ttn ttt ttg ttg atg ctg gna caa taa ttt agg tat tta
                                                                  193
Thr Gly Glu Xaa Phe Leu Leu Met Leu Xaa Gln Phe Arg Tyr Leu
         50
                             55
acc ata cat gct tta tac aac ata ttg tga gtt aca ata gcc ata aca
                                                                  241
Thr Ile His Ala Leu Tyr Asn Ile Leu Val Thr Ile Ala Ile Thr
                             70
cat tta tat tct ata taataacagt ag aat aat aga ata ttt ttt atg 292
His Leu Tyr Ser Ile
                                  Asn Asn Asn Arg Ile Phe Phe Met
         80
acc atttgtatct atacaatagt aaatagatta atacatataa gactatattc
                                                                  345
tttttgagag caacttaaag gagcggttat ggctttagtt acaaaagaag aagtacttca 405
ataccatagt gaaccccgac caggtaaact tgaagtattt tctataaaac catgtaaaac 465
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acaaaaagat cc
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His Ser Pro Leu Lys Arg Asn Val Xaa Met Glu Asp Val Gly Lys Ser
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Ala Val Phe Leu Ala Ser Asp Xaa Ser Ser Gly Val Thr Gly Glu Xaa
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Phe Leu Leu Met Leu Xaa Gln
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  Glu Leu Leu Val Phe Ser Gln Asn Arg Ser Gln Asn Ile Trp Leu Leu
                                        10
aca tta cct att ttt gtg tta ggt ata gca caa ggt ata tca ttt cct
                                                                    97
Thr Leu Pro Ile Phe Val Leu Gly Ile Ala Gln Gly Ile Ser Phe Pro
             20
tta qta aac agc cac att aca tca ctt qca cca aca tcc aac aga qct
Leu Val Asn Ser His Ile Thr Ser Leu Ala Pro Thr Ser Asn Arg Ala
         35
att gtt atg gct ata aac agt aca ttt atg agg tta agt cag agt att
                                                                   193
Ile Val Met Ala Ile Asn Ser Thr Phe Met Arg Leu Ser Gln Ser Ile
     50
                         55
tcg caa atg gtt ttt ggt att gga tgg tca ttt ttt ggt tgg cct ggt
Ser Gln Met Val Phe Gly Ile Gly Trp Ser Phe Phe Gly Trp Pro Gly
                     70
                                          75
cct ttt ata ttt ggt ctt ttt act tct att ata tta gcc ctc tta att
Pro Phe Ile Phe Gly Leu Phe Thr Ser Ile Ile Leu Ala Leu Leu Ile
                 85
                                      90
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Asn Asn Asn Arg Ile Phe Phe Met Thr

-	_			-	gta Val							_		-	337
					aaa Lys	_	_		_	_					385
					att Ile										433
tga		-	_		taa	-					-	-	-	_	481
		 _			tgg Trp		_	-		-			cc		525

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<212> PRT

<213> Lawsonia intracellularis

<400> 22

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Thr Leu Pro Ile Phe Val Leu Gly Ile Ala Gln Gly Ile Ser Phe Pro 20 25 30

Leu Val Asn Ser His Ile Thr Ser Leu Ala Pro Thr Ser Asn Arg Ala 35 40 45

Ile Val Met Ala Ile Asn Ser Thr Phe Met Arg Leu Ser Gln Ser Ile 50 55 60

Ser Gln Met Val Phe Gly Ile Gly Trp Ser Phe Phe Gly Trp Pro Gly 65 70 75 80

Pro Phe Ile Phe Gly Leu Phe Thr Ser Ile Ile Leu Ala Leu Leu Ile 85 90 95

Met Lys Tyr Phe Gln Asp Val Thr Gln Tyr His Leu Phe Leu Ile Ser 100 105 110

Ser Lys Phe Tyr Tyr 115 .

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Leu Ile Asp Ala Ile
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Ile Ser Ala Trp Asp Pro Gly
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ttttggatcct caacacaggg tatggattaa aacaacttta gctctaacag gagcatttta 180
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agaagacata ctacatatat ctactatacc tgtactacct ttctggaaag aatatacttc 600
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<210> 29

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<212> DNA

<213> Lawsonia intracellularis

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<211> 1082

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<213> Lawsonia intracellularis

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gtacatcaaa atattcttta cccaccttaa tacgaaaana aatnnttatn cnccncnatg 180
ggtggggntn aaatcctngc cccnttnccc tgttcnttta gggaaccccc naattccccn 240
ngttattcct ctgtttgaaa nttctggttn cccggccctn tnaccaanag cttgannncc 300
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<210> 32

<211> 477

<212> DNA

<213> Lawsonia intracellularis

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<210> 33

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<212> DNA

<213> Lawsonia intracellularis

<400> 33

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tccccatcat gataacatct cctaaaaaat aatttcatgg nggnaaaaat gttacctaca 420 catctctatt ttnaaagcaa aaaacccatg cccaanaaaa tttttgggcc naattaatat 480 acttaatcta ataaactttt ttgggtaatn aaaaaaaatt aatttttaa acttggtttn 540 accaaccttt tctccttact ttttaacc 568

<210> 34

<211> 477

<212> DNA

<213> Lawsonia intracellularis

<400> 34

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